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Sequence Listing was accepted.

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217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=9; day=16; hr=11; min=44; sec=31; ms=520;]

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Application No: 10815480

Version No: 2.0

Input Set:**Output Set:****Started:** 2008-08-22 13:43:19.010**Finished:** 2008-08-22 13:43:26.690**Elapsed:** 0 hr(s) 0 min(s) 7 sec(s) 680 ms**Total Warnings:** 379**Total Errors:** 0**No. of SeqIDs Defined:** 919**Actual SeqID Count:** 919

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Started: 2008-08-22 13:43:19.010
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Actual SeqID Count: 919

Error code

Error Description

This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Young, Karen K. Y.

<120> Compositions and Methods for Detecting Certain
Flaviviruses, Including Members of the Japanese
Encephalitis Virus Serogroup

<130> 022101-000230US

<140> 10815480

<141> 2004-03-31

<150> US 60/459,491

<151> 2003-03-31

<150> US 60/552,454

<151> 2004-03-12

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<223> Example Primer 1, Japanese encephalitis virus
serogroup amplification primer

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 virus serogroup, KY1129

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nucleic acid

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<223> probe for detecting Japanese encephalitis virus
serogroup members

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<223> probe for detecting yellow fever virus

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<211> 28

<212> DNA

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<223> probe for detecting Montana myotis
leukencephalitis virus

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<211> 28

<212> DNA

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<223> probe for detecting Modoc virus

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<211> 28

<212> DNA

<213> Artificial Sequence

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<223> Example Probe 1, Japanese encephalitis virus
serogroup amplification primer

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<211> 418

<212> DNA

<213> St. Louis encephalitis virus

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<223> 3' untranslated region of the genome of St. Louis
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gtaggctgga gaggacgcaa aagtcagacc agaaatgcca cctgaaagca tgctaaaggt 180
gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
aaccatggag tgcgtgacca tggcgtaagg actagagggt agaggagacc ccgctgcaac 300
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gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
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gtaggctgga gaggacgcaa aagtcagacc agaaatgcca cctgaaagca tgctaaaggt 180
gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
aaccatggag tgcgtgacca tggcgtaagg actagagggt agaggagacc ccgctgcaac 300
tcggcaaggc ccaaaccgc tcgaagctgt agagatgggg gaaggactag aggttagagg 360
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<223> 3' untranslated region of the genome of St. Louis
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gtaggctgga gaggacgcaa aagtcagacc agaaatgcca cctgaaagca tgctaaaggt 180
gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
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gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
aaccatggag tgcgtgacca tggcgtaagg actagaggtt agaggagacc ccgctgtaac 300
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gtaggctgga gaggacgcac aagtcagacc agaaatgcca cctgaaagca tgctaaaggt 180
gctgtctgta catgccccag gaggactggg ttaacaaagc ttaacagccc cagcggccca 240
aaccatggag agcgtgacca tggcgtaagg actagaggtt agaggagacc ccgctgtaac 300
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gtaggctgga gaggacgcac aagtcagacc agaaatgcca cctgaaagca tgctaaaggt 180
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aaccatggag tgcgtgacca tggcgtaagg actagagggt agaggagacc ccgctgtaac 300
ttggcaaggc ccaaaccgcg tcaaagctgt agagacgggg gaaggactag aggttagagg 360
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<223> 3' untranslated region of the genome of St. Louis
encephalitis virus (SLEV) isolate Kern217

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gaaagcatgc taaaggtgct gtctgtacat gcccaggag gactgggtta acaaagctta 180
acagccccag cggcccaaac catggagtgc gtgacctgg cgtaaggact agaggttaga 240
ggagaccccg ctgtaacttg gcaaggccca aaccgctca aagctgtaga gacgggggaa 300
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aaccatggag tgcgtgacca tggcgtaagg actagagggt agaggagacc ccgctgtaat 300
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ggagaccccg ctgcaacttg gcaaggccca aaccgctcg aagctgtaga gacgggggaa 300
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